

Figure 1 SEQ ID NO:1

-Sequence ID-1: human interferon alpha 1 (GenBank 13128950)

MASPFALLMVLVVLSCKSSCSLGCDLPETHSLDNRRTLMLLAQMSRISPSSCLMDRHDFGF PQEEFDGNQFQKAPAISVLHELIQQIFNLFTTKDSSAAWDEDLLDKFCTELYQQLNDLEACV MQEERVGETPLMNADSILAVKKYFRRITLYLTEKKYSPCAWEVVRAEIMRSLSLSTNLQERL RRKE

SEQ IDNO: 2

Sequence ID-2: human interferon alpha-2a (GenBank 2781226)

CDLPQTHSLGSRRTLMLLAQMRKISLFSCLKDRHDFGFPQEEFGNQFQKAETIPVLHEMIQ QIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLEACVIQGVGVTETPLMKEDSILAVRKYF QRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE

SEQ LDNO: 3

Sequence ID-3: human interferon alpha-2b (GenBank 30171279)

MCDLPQTHSLGSRRTLMLLAQMRRISLFSCLKDRHDFGFPQEEFGNQFQKAETIPVLHEMI QQIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLEACVIQGVGVTETPLMKEDSILAVRKY FQRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE

SEQ ID NO: 4

Sequence ID-4: human interferon alpha 4 (GenBank 10835103)

MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISHFSCLKDRHDFGFP EEEFDGHQFQKAQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLEACVI QEVGVEETPLMNEDSILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTNLQKRLR RKD

SER ID NO:5

Sequence ID 5: human interferon alpha 5 (GenBank 4504597)

MALPFVLLMALVVLNCKSICSLGCDLPQTHSLSNRRTLMIMAQMGRISPFSCLKDRHDFGFP QEEFDGNQFQKAQAISVLHEMIQQTFNLFSTKDSSATWDETLLDKFYTELYQQLNDLEACM MQEVGVEDTPLMNVDSILTVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSLSANLQERL RRKE

SER IDNO 6

Sequence ID 6: human interferon alpha 6 (GenBank 11128015)

MALPFALLMALVVLSCKSSCSLDCDLPQTHSLGHRRTMMLLAQMRRISLFSCLKDRHDFRF PQEEFDGNQFQKAEAISVLHEVIQQTFNLFSTKDSSVAWDERLLDKLYTELYQQLNDLEAC VMQEVWVGGTPLMNEDSILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSSSRNLQE RLRRKE

SEQIDNO: 7

Sequence ID-7: human interferon alpha 7 (GenBank 10800142)

MARSFSLLMAVLVLSYKSICSLGCDLPQTHSLRNRRALILLAQMGRISPFSCLKDRHEFRFP EEEFDGHQFQKTQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLEACVI QEVGVEETPLMNEDFILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSFSTNLKKGLR RKD

SPRIDNO! 8

-Sequence-ID-8: human interferon alpha 8 (GenBank 4504599)

MALTFYLMVALVVLSYKSFSSLGCDLPQTHSLGNRRALILLAQMRRISPFSCLKDRHDFEFP QEEFDDKQFQKAQAISVLHEMIQQTFNLFSTKDSSAALDETLLDEFYIELDQQLNDLEVLCD QEVGVIESPLMYEDSILAVRKYFQRITLYLTEKKYSSCAWEVVRAEIMRSFSLSINLQKRLKS KE

FIG. _ 1A

SEQ ID NO:9

Sequence ID 9: human interferon alpha 10 (GenBank 4504589)

MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLGQMGRISPFSCLKDRHDFRIPQ EEFDGNQFQKAQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLEACVIQ EVGVEETPLMNEDSILAVRKYFQRITLYLIERKYSPCAWEVVRAEIMRSLSFSTNLQKRLRRK D

SEQ IDNO: 10

Sequence ID-10: human interferon alpha 13 (GenBank 13128966)

MASPFALLMALVVLSCKSSCSLGCDLPETHSLDNRRTLMLLAQMSRISPSSCLMDRHDFGF PQEEFDGNQFQKAPAISVLHELIQQIFNLFTTKDSSAAWDEDLLDKFCTELYQQLNDLEACV MQEERVGETPLMNADSILAVKKYFRRITLYLTEKKYSPCAWEVVRAEIMRSLSLSTNLQERL RRKE

SEQ IDNO:1

~Sequence ID 1'1: human interferon alpha 14 (GenBank 4504591)

MALPFALMMALVVLSCKSSCSLGCNLSQTHSLNNRRTLMLMAQMRRISPFSCLKDRHDFE FPQEEFDGNQFQKAQAISVLHEMMQQTFNLFSTKNSSAAWDETLLEKFYIELFQQMNDLEA CVIQEVGVEETPLMNEDSILAVKKYFQRITLYLMEKKYSPCAWEVVRAEIMRSFSFSTNLQK RLRRKD

SEQ 10 NO: 12

Sequence ID 12: human interferon alpha 16 (GenBank 4504593)

MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISHFSCLKDRYDFGFP QEVFDGNQFQKAQAISAFHEMIQQTFNLFSTKDSSAAWDETLLDKFYIELFQQLNDLEACVT QEVGVEEIALMNEDSILAVRKYFQRITLYLMGKKYSPCAWEVVRAEIMRSFSFSTNLQKGLR RKD

SEQ 10 NO: 13

Sequence ID 13: human interferon alpha 17 (GenBank 10880985)

MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISPFSCLKDRHDFGLP QEEFDGNQFQKTQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNNLEACVI QEVGMEETPLMNEDSILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTNLQKILR RKD

SPQ 10 NO: 14

Sequence ID-14: human interferon alpha 21 (4504595)

MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISPFSCLKDRHDFGFP QEEFDGNQFQKAQAISVLHEMIQQTFNLFSTKDSSATWEQSLLEKFSTELNQQLNDMEACV IQEVGVEETPLMNVDSILAVKKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSLSKIFQERLR RKE

SERID NO:15

Sequence 1D-15: human interferon beta (GenBank 124469), signal peptide deleted MSYNLLGFLQRSSNFQCQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDFTRGKLMSSLHLKR YYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SEQIDMO:16

Sequence ID 16: human interferon kappa (GenBank 14488028)

MSTKPDMIQKCLWLEILMGIFIAGTLSLDCNLLNVHLRRVTWQNLRHLSSMSNSFPVECLRE NIAFELPQEFLQYTQPMKRDIKKAFYEMSLQAFNIFSQHTFKYWKERHLKQIQIGLDQQAEY

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belongs on [FIB

LNQCLEEDENENEDMKEMKENEMKPSEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEI VRVEIRRCLYYFYKFTALFRRK

980 1D NO:17

-Sequence ID-17: human interferon tau (GenBank 28882045)

MIIKHFFGTVLVLLASTTIFSLDLKLIIFQQRQVNQESLKLLNKLQTLSIQQCLPHRKNFLLPQK SLSPQQYQKGHTLAILHEMLQQIFSLFRANISLDGWEENHTEKFLIQLHQQLEYLEALMGLE AEKLSGTLGSDNLRLQVKMYFRRIHDYLENQDYSTCAWAIVQVEISRCLFFVFSLTEKLSKQ GRPLNDMKQELTTEFRSPR

SEQ 1D NO:18

Sequence ID-18: human interferon omega (GenBank 4504605)

MALLFPLLAALVMTSYSPVGSLGCDLPQNHGLLSRNTLVLLHQMRRISPFLCLKDRRDFRFP QEMVKGSQLQKAHVMSVLHEMLQQIFSLFHTERSSAAWNMTLLDQLHTGLHQQLQHLETC LLQVVGEGESAGAISSPALTLRRYFQGIRVYLKEKKYSDCAWEVVRMEIMKSLFLSTNMQE RLRSKDRDLGSS

SERIDNO: 19

Sequence ID-19: interferon beta variant #2 L5Q in C17S background

MSYNQLGFLQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDFTRGKLMSSLHLKR YYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SOLIDNO: 20

Sequence-ID-20:-interferon beta variant #7 L5Q/F8E in C17S background MSYNQLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDFTRGKLMSSLHLKR YYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SEQ 10 NO: 21

Sequence-ID-21: interferon beta variant #15 L5Q/F8E/F111N in C17S background MSYNQLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDNTRGKLMSSLHLK RYYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SER 10 NO. 22

Sequence ID 22: interferon beta variant #23 L5Q/F8E/L116E in C17S background MSYNQLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDFTRGKEMSSLHLK RYYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SEQ 10 NO:23

Sequence ID 23: interferon beta variant #36 F8E/F111N/L116E in C17S background MSYNLLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDNTRGKEMSSLHLK RYYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

SERIONO: 24

Sequence ID 24: interferon beta variant #39 L5Q/F8E/F111N/L116E in C17S background MSYNQLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDNTRGKEMSSLHLK RYYGRILHYLKAKEYSHCAWTIVRVEILRNFYFINRLTGYLRN

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4113

SERIDNO: 25

Sequence ID 25: interferon beta variant #64 L5Q/F8E/L47K/F111N/L116E/L120R in C17S background

MSYNQLGELQRSSNFQSQKLLWQLNGRLEYCLKDRMNFDIPEEIKQKQQFQKEDAALTIYE MLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLKTVLEEKLEKEDNTRGKEMSSRHLK RYYGRILHYLKAKEYSHCAWTIVRVEI LRNFYFINRLTGYLRN

SEQ 10NO:26

Sequence ID 26: interferon kappa variant #4_G7 V8N/W15R/Y48Q/M52N/F76S/Y78A/I89T LDCNLLNNHLRRVTRQNLRHLSSMSNSFPVECLRENIAFELPQEFLQQTQPNKRDIKKAFYE MSLQAFNIFSQHTSKAWKERHLKQIQTGLDQQAEYLNQCLEEDENENEDMKEMKENEMKP SEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRVEIRRCLYYFYKFTALFRRK

SEQ ID NO.27

Sequence ID 27: interferon kappa variant #46_E2 W15R/I37N/Y48Q/M52N/F76S/Y78A/I89T/ Y97D/V161A/C166A/Y168S/Y171T LDCNLLNVHLRRVTRQNLRHLSSMSNSFPVECLRENNAFELPQEFLQQTQPNKRDIKKAFY EMSLQAFNIFSQHTSKAWKERHLKQIQTGLDQQAEDLNQCLEEDENENEDMKEMKENEMK PSEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRAEIRRALSYFTKFTALFRRK

SEQ ID NIO:28

Sequence ID-28: interferon kappa variant #47_C4 W15R/F76S/Y78A

LDCNLLNVHLRRVTRQNLRHLSSMSNSFPVECLRENIAFELPQEFLQYTQPMKRDIKKAFYE MSLQAFNIFSQHTSKAWKERHLKQIQIGLDQQAEYLNQCLEEDENENEDMKEMKENEMKP SEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRVEIRRCLYYFYKFTALFRRK

SER ID NO:29

Sequence ID-29: interferon kappa variant #23_C10 I37N/Y48Q/M52N/F76S/Y78A/Y97D LDCNLLNVHLRRVTWQNLRHLSSMSNSFPVECLRENNAFELPQEFLQQTQPNKRDIKKAFY EMSLQAFNIFSQHTSKAWKERHLKQIQIGLDQQAEDLNQCLEEDENENEDMKEMKENEMK PSEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRVEIRRCLYYFYKFTALFRRK

SEQ 10 NO: 30

Sequence ID 30: interferon kappa variant #40_A10 W15R/I37N/F76S/Y78A LDCNLLNVHLRRVTRQNLRHLSSMSNSFPVECLRENNAFELPQEFLQYTQPMKRDIKKAFY EMSLQAFNIFSQHTSKAWKERHLKQIQIGLDQQAEYLNQCLEEDENENEDMKEMKENEMK PSEARVPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRVEIRRCLYYFYKFTALFRRK

F161. - 1D

Figure 2

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CDLPETHSLDNRRTLMLLAQMSRISPSSCLMDRHDFGFPQEEFDGNQFQKAPAISVLHEL 83
     1: 24
32 2a: 2
            CDLPQTHSLGSRRTLMLLAQMRKISLFSCLKDRHDFGFPOEEF-GNOFOKAETIPVLHEM 60
33 2b: 2
            CDLPQTHSLGSRRTLMLLAQMRRISLFSCLKDRHDFGFPQEEF-GNQFQKAETIPVLHEM 60
34 4: 24
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 36 6: 24
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 37 7: 24
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 388: 24
            CDLPQTHSLGNRRALILLAQMRRISPFSCLKDRHDFEFPQEEFDDKQFQKAQAISVLHEM 83
 39 10: 24
            CDLPQTHSLGNRRALILLGQMGRISPFSCLKDRHDFRIPQEEFDGNQFQKAQAISVLHEM 83
 40 13: 24
            CDLPETHSLDNRRTLMLLAQMSRISPSSCLMDRHDFGFPQEEFDGNQFQKAPAISVLHEL 83
 41 14: 24
            CNLSQTHSLNNRRTLMLMAQMRRISPFSCLKDRHDFEFPQEEFDGNOFOKAOAISVLHEM 83
 4216: 24
            CDLPQTHSLGNRRALILLAQMGRISHFSCLKDRYDFGFPQEVFDGNQFQKAQAISAFHEM 83
 43 17: 24
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 44 21: 24
            CDLPQTHSLGNRRALILLAQMGRISPFSCLKDRHDFGFPOEEFDGNOFOKAOAISVLHEM 83
31
     1: 84
            IQQIFNLFTTKDSSAAWDEDLLDKFCTELYQQLNDLEACVMQEERVGETPLMNADSILAV 143
32 2a: 61
            IQQIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLEACVIQGVGVTETPLMKEDSILAV 120
33 2b: 61
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34 4: 84
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 37 7: 84
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38 8: 84
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4013: 84
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4/ 14: 84
           MQQTFNLFSTKNSSAAWDETLLEKFYIELFQQMNDLEACVIQEVGVEETPLMNEDSILAV 143
42 16: 84
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43 17: 84
           IQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNNLEACVIQEVGMEETPLMNEDSILAV 143
4421: 84
           IQQTFNLFSTKDSSATWEQSLLEKFSTELNQQLNDMEACVIQEVGVEETPLMNVDSILAV 143
    1: 144 KKYFRRITLYLTEKKYSPCAWEVVRAEIMRSLSLSTNLQERLRRKE 189
32 2a: 121 RKYFQRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE 166
33 2b: 121 RKYFQRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE 166
34 4: 144 RKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTNLQKRLRRKD 189
35 5: 144 RKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSLSANLQERLRRKE 189
36 6: 144 RKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSSSRNLQERLRRKE 189
37 7: 144 RKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSFSTNLKKGLRRKD 189
38 8: 144 RKYFQRITLYLTEKKYSSCAWEVVRAEIMRSFSLSINLQKRLKSKE 189
39 10: 144 RKYFQRITLYLIERKYSPCAWEVVRAEIMRSLSFSTNLQKRLRRKD 189
40 13: 144 KKYFRRITLYLTEKKYSPCAWEVVRAEIMRSLSLSTNLQERLRRKE 189
4) 14: 144 KKYFQRITLYLMEKKYSPCAWEVVRAEIMRSFSFSTNLQKRLRRKD 189
4216: 144 RKYFQRITLYLMGKKYSPCAWEVVRAEIMRSFSFSTNLQKGLRRKD 189
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4317: 144 RKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTNLOKILRRKD 189

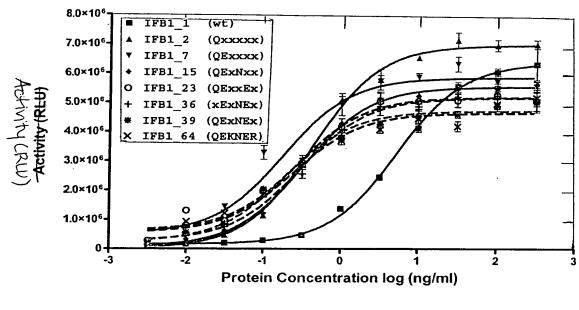
47 21: [144 KKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSLSKIFQERLRRKE 189] - Fort Size charged to smaller size

Figure 3

| 45 IFNK: 46 1AU1: 47 1B5L: | MSYNLLGFLQRSSNFQCQKLLWQLNGRLEY-CLKDRMNFDIPEEIKQLQQ |
|----------------------------------|--|
| 48 11TF: | 2 |
| | ${\tt pmkrdikkafyemslqafnifsqhtfkywkerhkqiqigldqqaeyln}$ |
| 46 1AU1: | FQKEDAALTIYEMLQNIFAIFRQDSSSTGWNETIVENLLANVYHQINHLK |
| 47 1B5L: | LQKDQAFPVLYEMLQQSFNLFYTEHSSAAWDTTLLEQLCTGLQQQLDHLD |
| 48 11TF: | FQKAETIPVLHEMIQQIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLE |
| | qcleedenenedmkemkenemkpsearvpqlsslelrryfhridnflkek |
| | TVLEEKLEKEDFTRGKLMSSLHLKRYYGRILHYLKAK |
| 471B5L: | TCRG MDPIVTVKKYFQGIYDYLQEK |
| 4811TF: | ACVIQGVGVTETPLMKEDSILAVRKYFQRITLYLKEK |
| | kysdcaweivrveirrclyyfykftalfrrk |
| 40 1AU1: | EYSHCAWTIVRVEILRNFYFINRLTGYLRN |
| 471B5L: | GYSDCAWEIVRVEMMRALTVSTTLQKRLTK |
| | KYSPCAWEVVRAEIMRSFSLSTNLQESLRSKE |

FIG.-3

Figure 4



F16.-4

716._4 has been enlarged to fit page, oriented as Landscape

Figure_5-

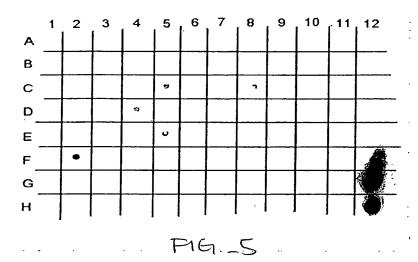
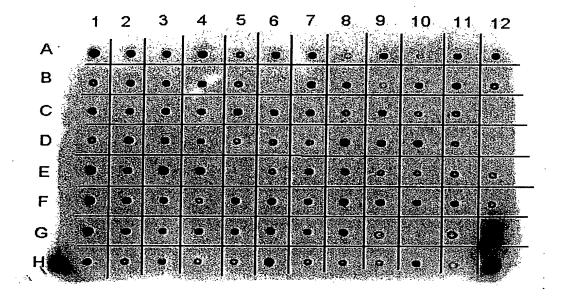


FIG._4 Appears here

F161.-6

which

Figure 6



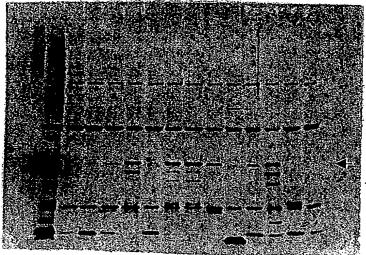
P16.- 6

Fig. 16 appears on page 8/13 with Fig. -5

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-Figure-7-

IK1 RETEST PLATE #1 WESTERN #3



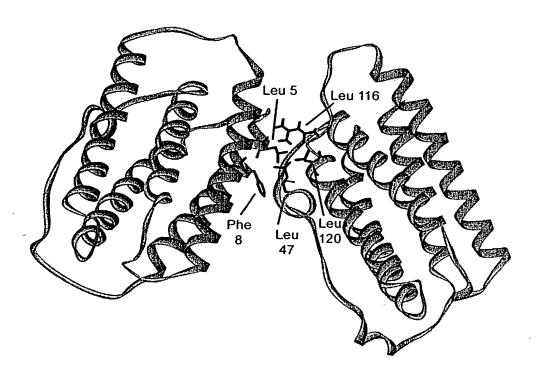
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15

P161._7

FIG._8 appears here

F19.-8

Figure-8-



F16._8

FIG. 2 appears on page 9/13 below FIG. 7. Figure 2

-5-

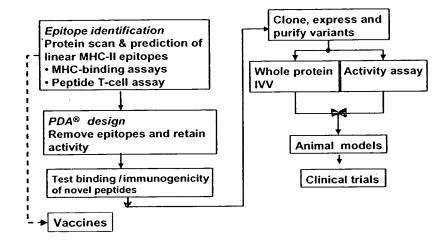
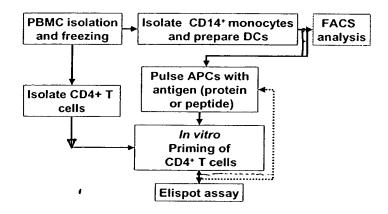


FIG. -9

716._10 appears

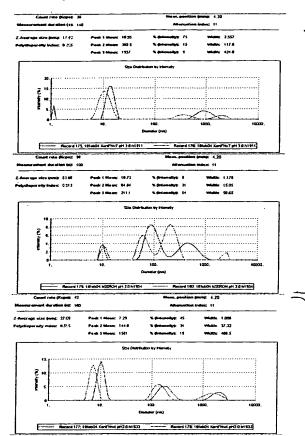
Figure & 10



PIG ._ 10

F161.-10 appears on page 10/13 below F161.-9 and is enlarged

Figure



Now FIG. 11A and is enlarged

Now Pig._IIB and is enlarged

Now FIG._11C - enlarged and appears on page 12/13.

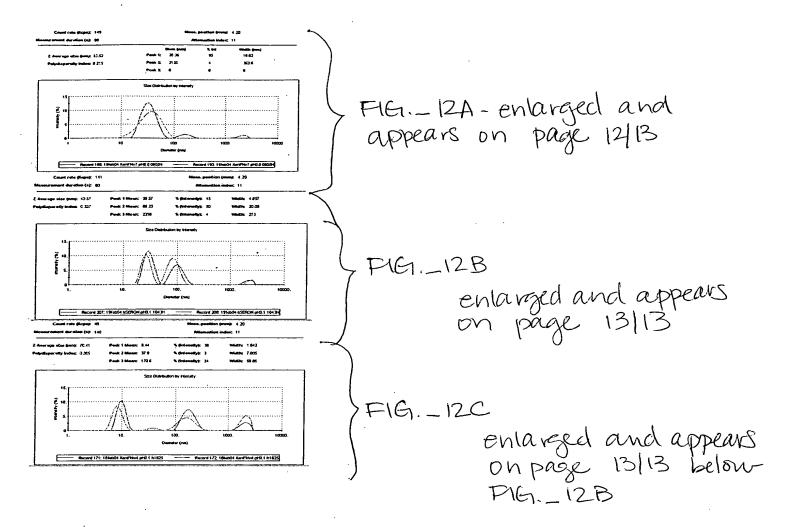


Figure 1M 12